

↓

A)	<u>MRGATRVSIMLLLVTVSDC</u>	AVITGA
B)	<u>MRSLCCAPLLLLLLLPLLLTPRAGDA</u>	AVITGA
C)	<u>MKCFAIQIVLLLLVIAFSHG</u>	AVITGA
D)		AVITGA

CERDVQCGAGTCCAI SLWLRGLRMCTPLGREGEECHPG
CDKDSQCGGGMCCAVSIWVKSIRICTPMGKLGDSCHPL
CDKDVQCGSGTCCAASAWSRNIRFCIPLGNSGEDCHPA
CERDLQCGKGTCCAVSLWIKSVRVCTPVGTSGEDCHPA
* * ** * *

SHKVPFFRKRKHHTCPCLPNLLCSRFPDGRYRCSMDLKNINF
TRKVPFFGRMHHTCPCLPGLACLRTSFNRFICLAQK
SHKVPYDGKRLSSLCPCCKSGLTCSK.SGEKFKCS
SHKIPFSGQRMHTCPCAPNLACVQTSPKKFKCLSKS
 * * * *

FIGURE 1

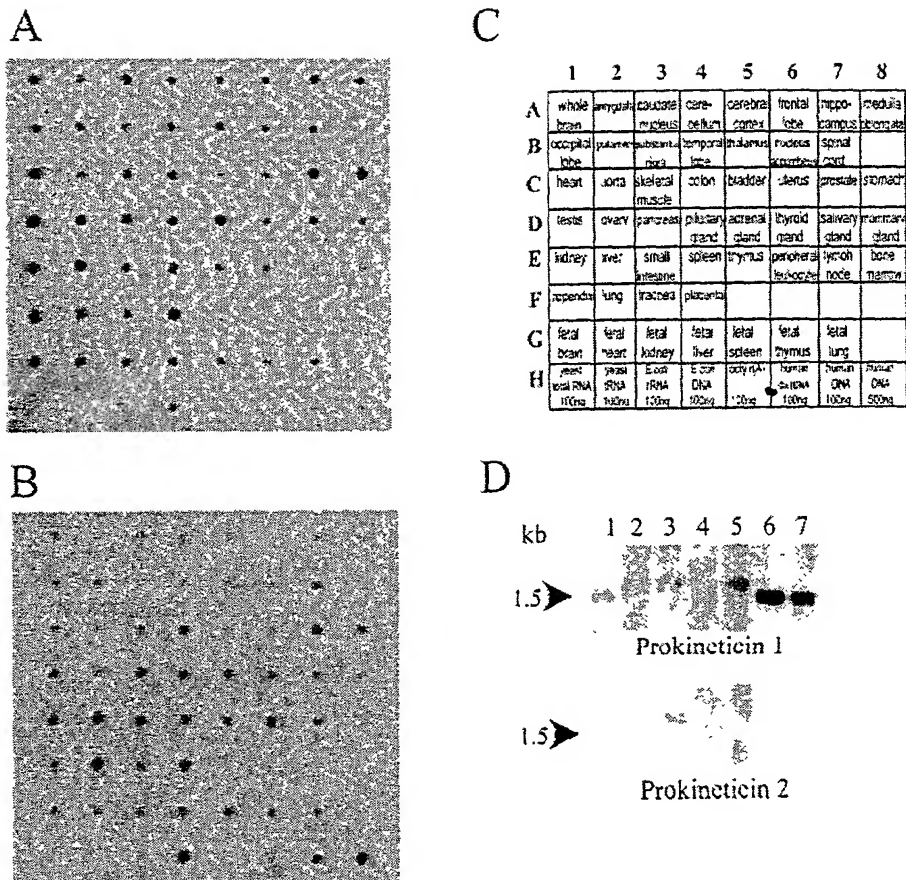
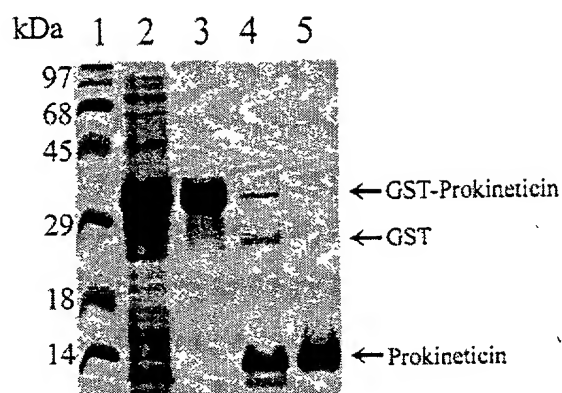
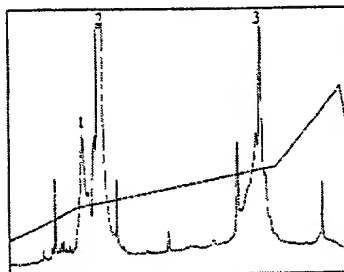


FIGURE 2

A



B



C

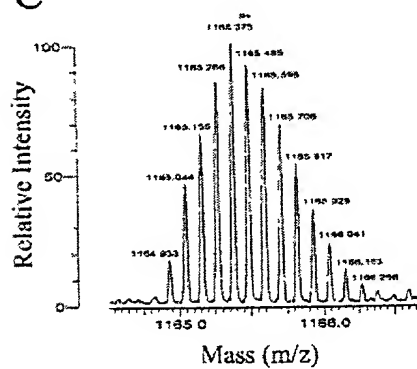


FIGURE 3

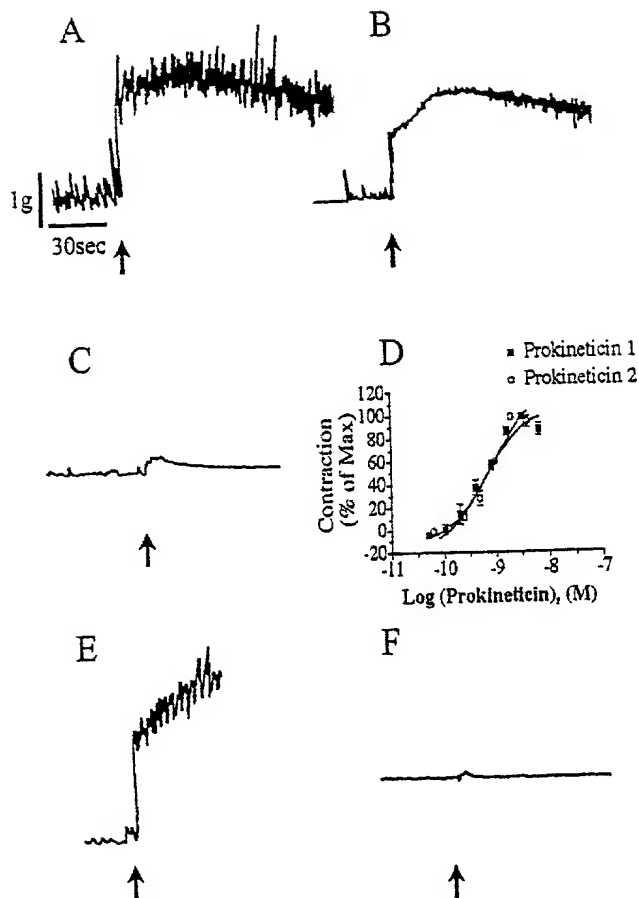
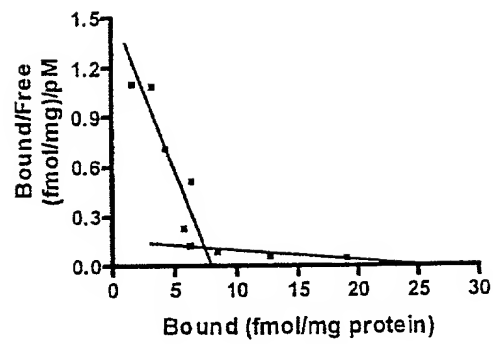


FIGURE 4

A



B

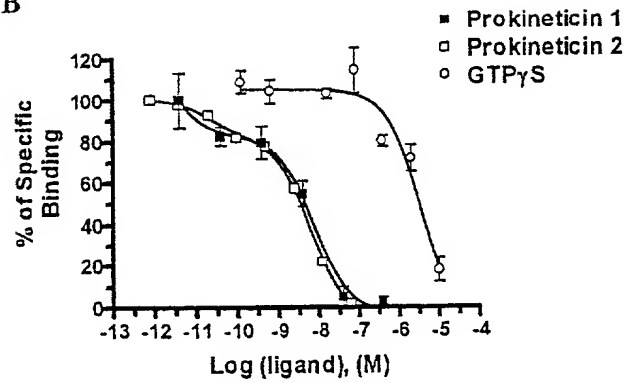


FIGURE 5

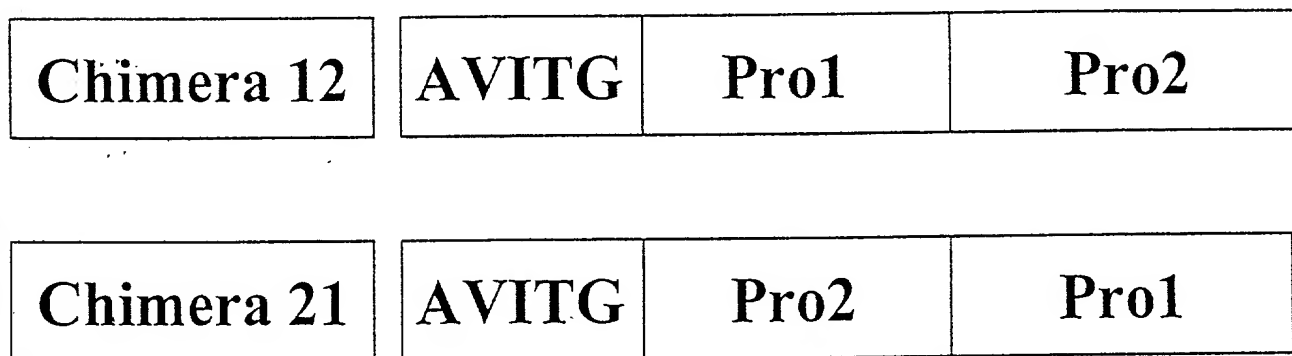
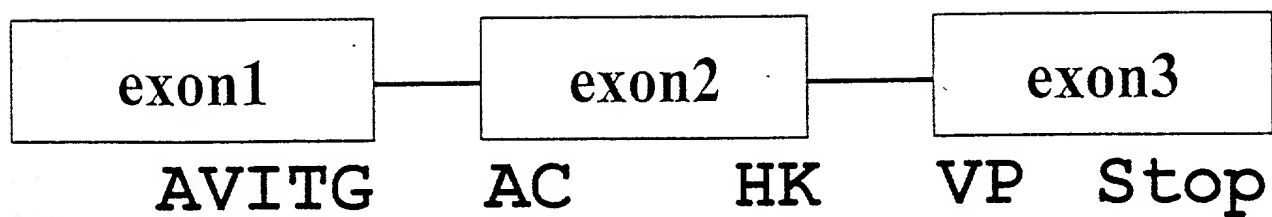
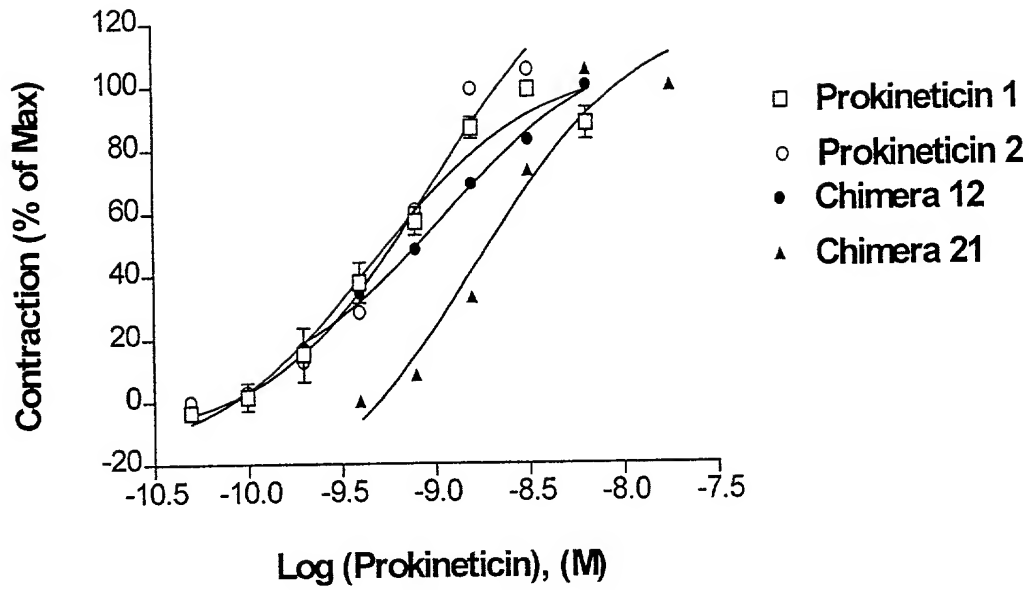


FIGURE 6

A



B

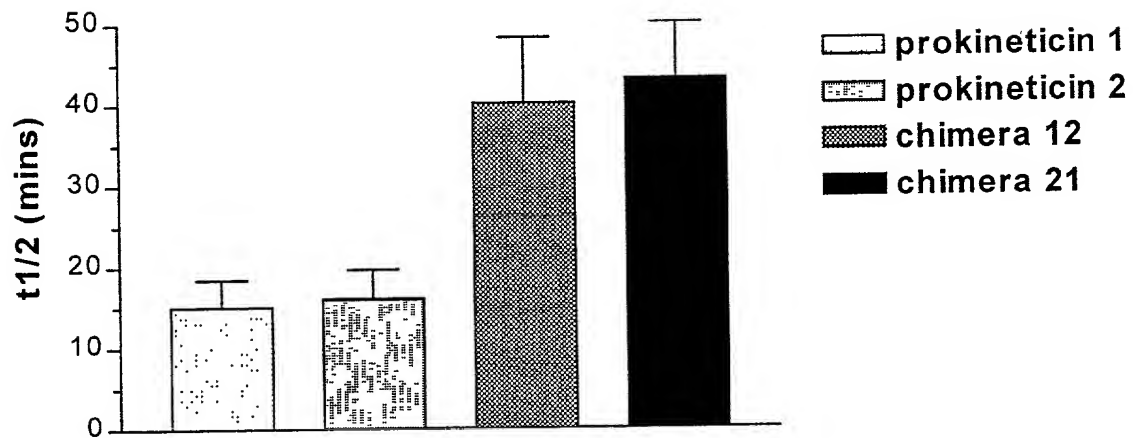


FIGURE 7

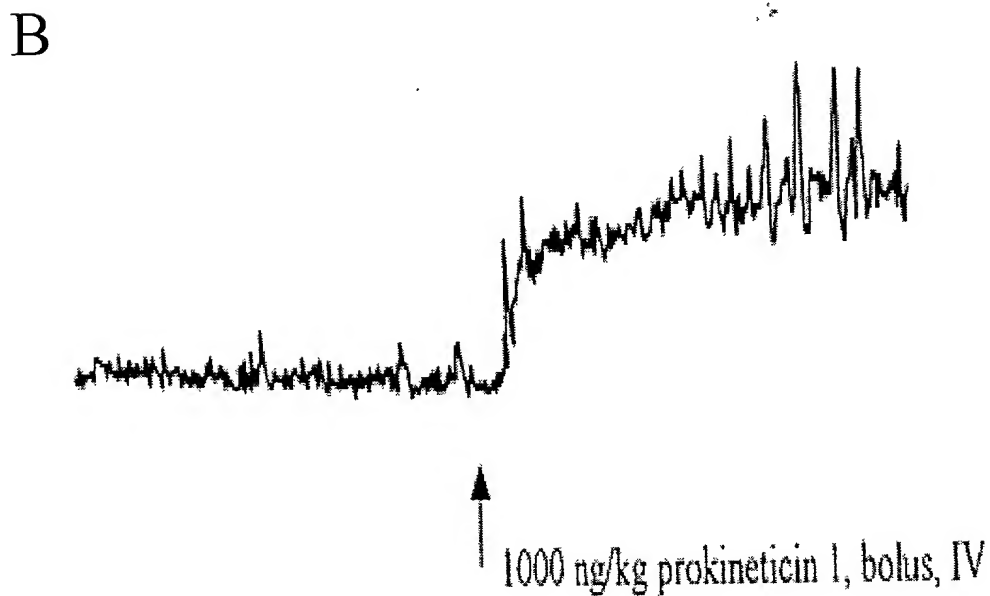
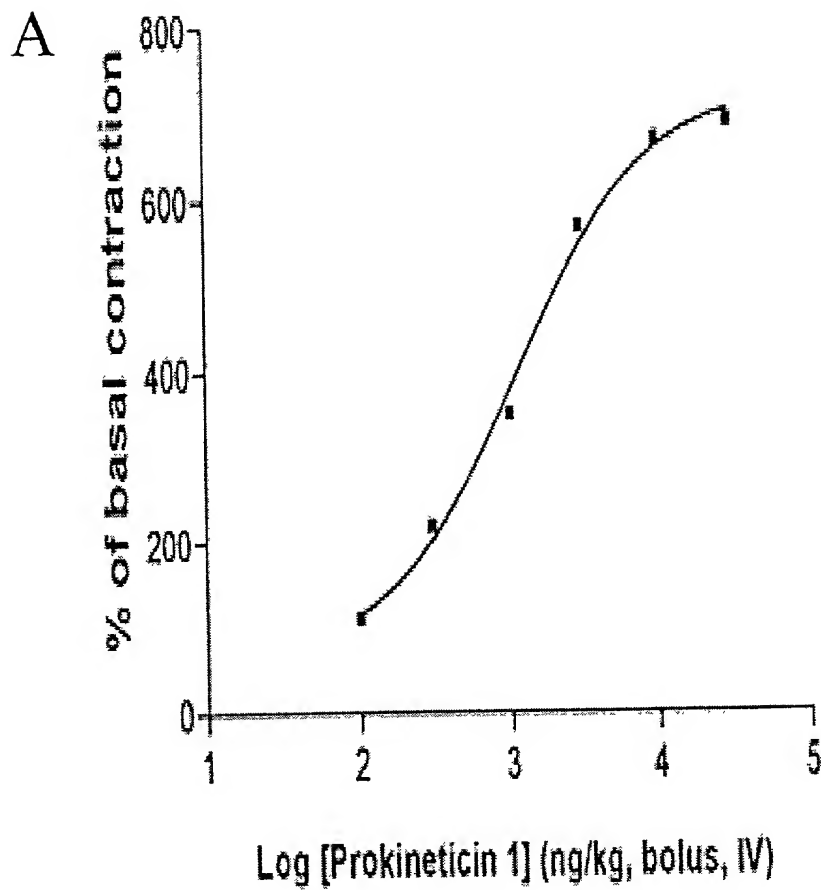


FIGURE 8

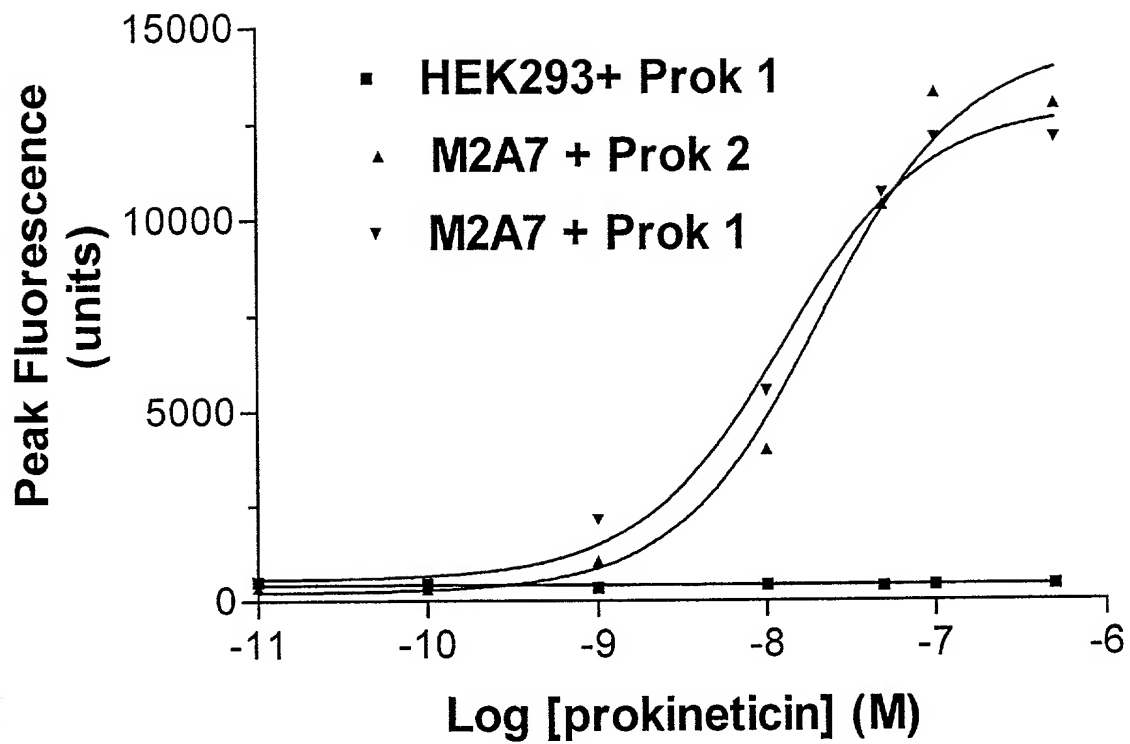


FIGURE 9